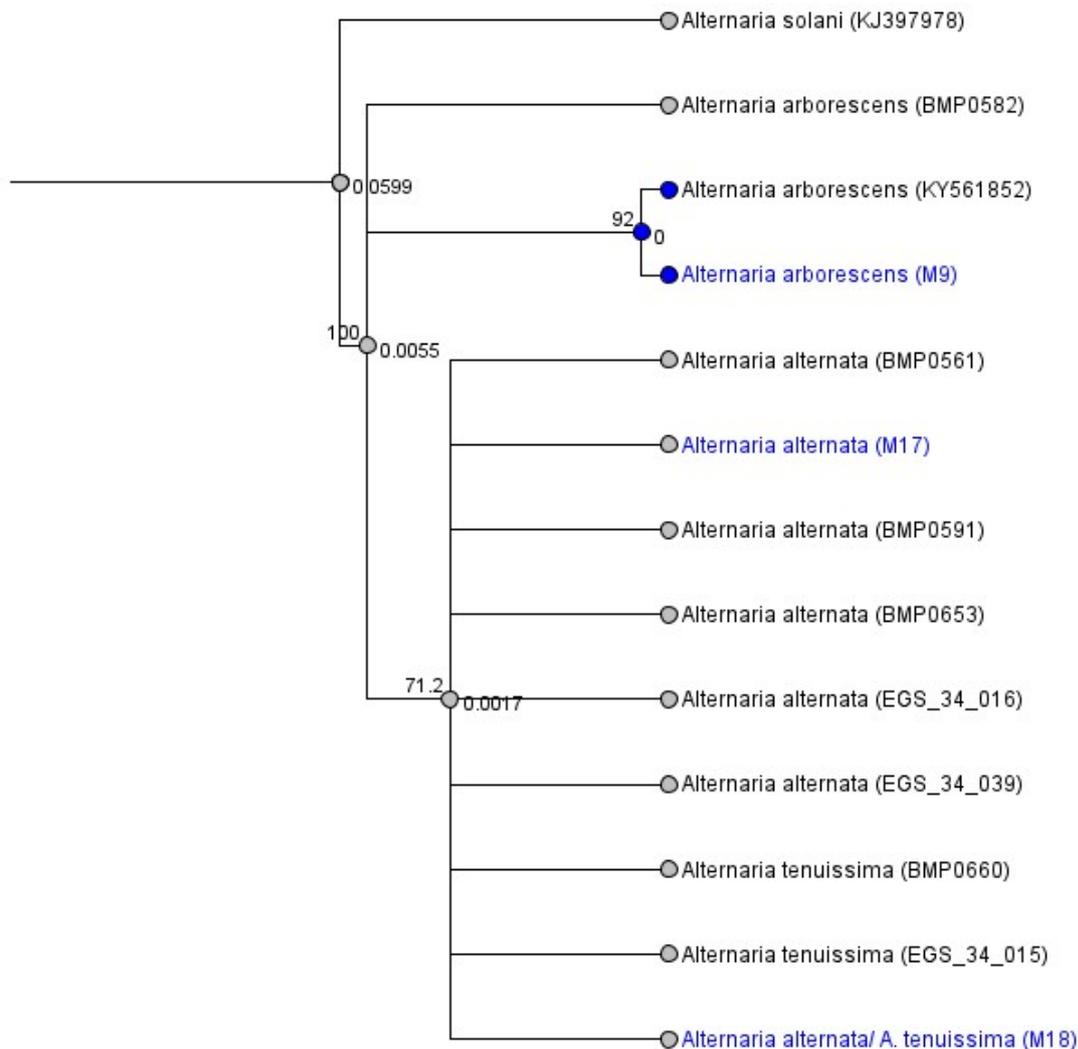


Article

# Species of the genera *Neopestalotiopsis* and *Alternaria* as dominant pathogen species attacking mastic trees (*Pistacia lentiscus* var. Chia)



**Figure S2.** The phylogenetic tree was generated by Geneious Prime® 2022.1.1 (Biomatters Ltd, New Zealand) software analyzing the data set of ITS gene as downloaded from the National Center for Biotechnology Information (NCBI) database. The Tamura – Nei genetic distance model was used, and the tree was constructed according to UPGMA method. *Alternaria solani* was used as the outgroup and the strains isolated within this study are written in blue. The accession numbers of all strains used are the following: *Alternaria solani* (KJ397978), *Alternaria arborescens* (BMP0582), *Alternaria arborescens* (KY561852), *Alternaria alternata* (BMP0561), *Alternaria alternata* (BMP0591), *Alternaria alternata* (BMP0653), *Alternaria alternata* (EGS\_34\_016), *Alternaria alternata* (EGS\_34\_039), *Alternaria alternata* (BMP0660), *Alternaria tenuissima* (EGS\_34\_015).